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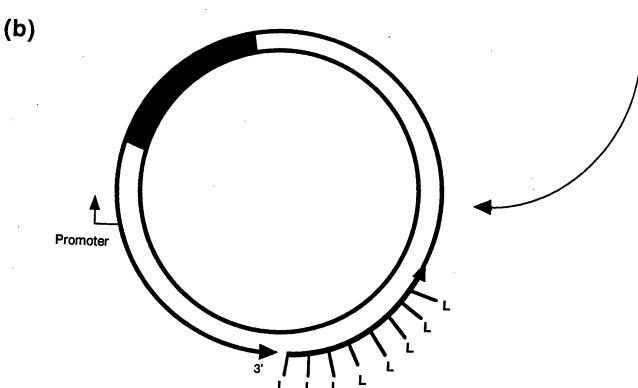


Figure I
Attachment of Ligands Through Primer Region

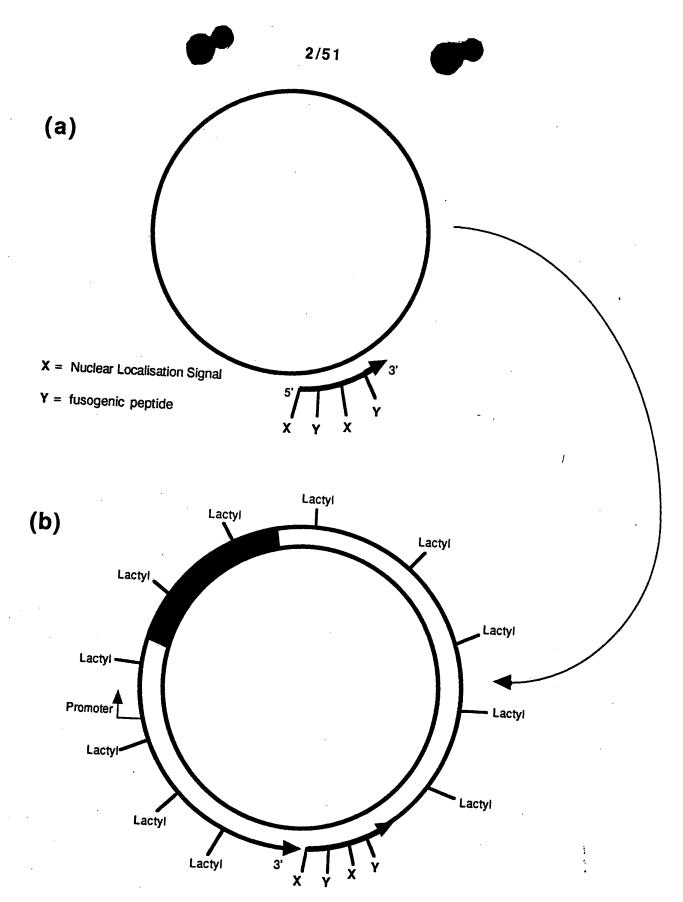


Figure 2
Attachment of Ligands by Incorporation of Modified Nucleotide Precursors

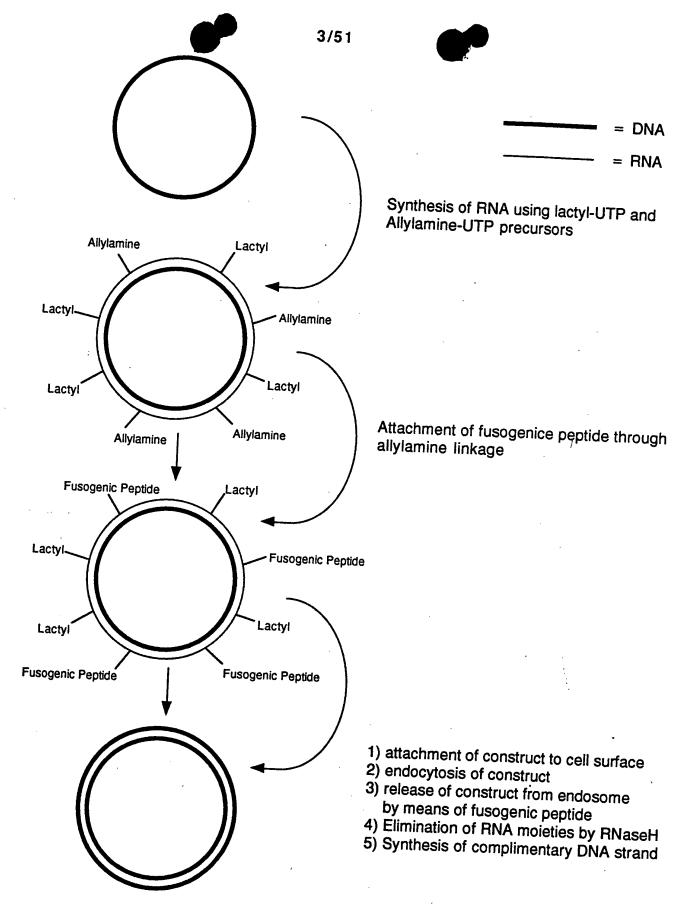


Figure 3
Incorporation of Ligands through Modified Ribonucleotides

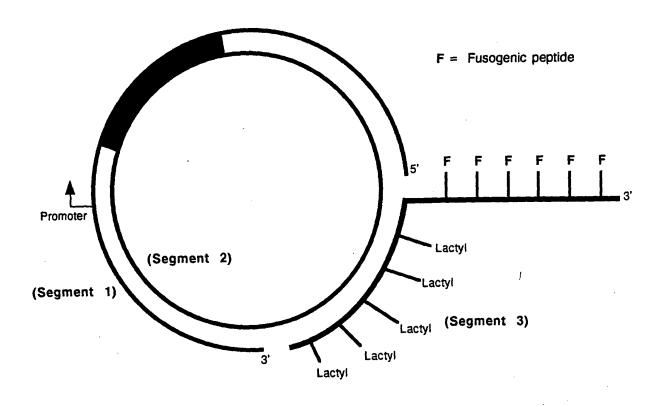


Figure 4

Attachment of Ligands through a 3' tail

Preparation of Gapped Circle

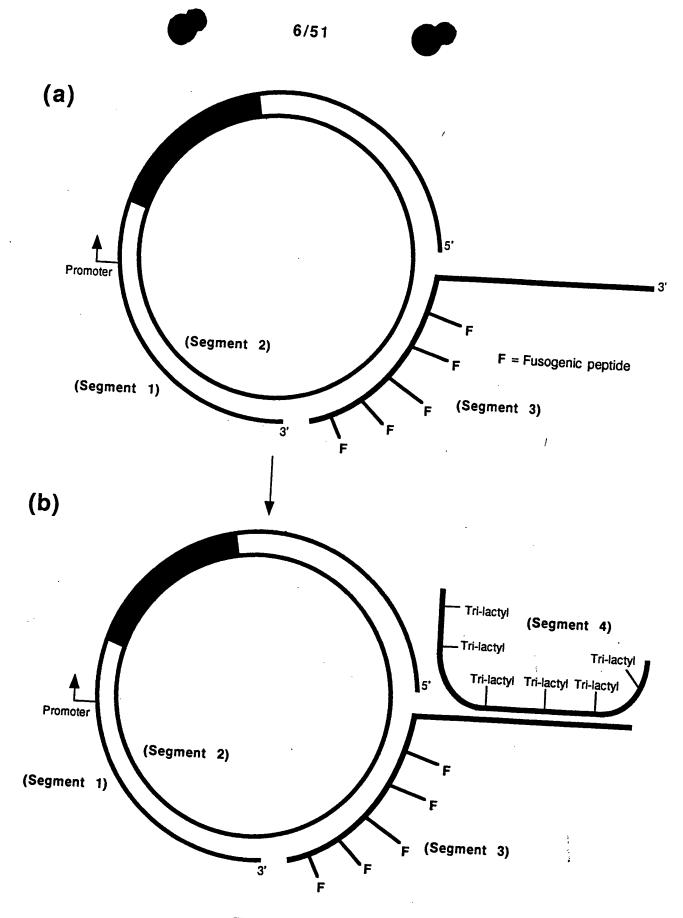


Figure 6
Attachment of Ligands through hybridization to a 3' tail

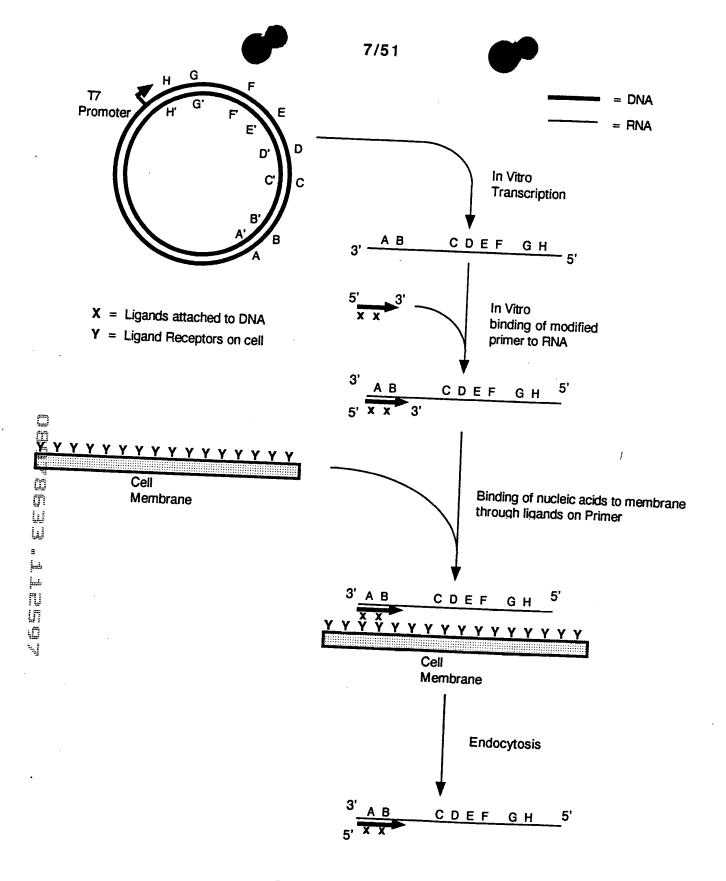


Figure 7
RNA with Ligands on Primer

(Continued in Figure 8)





Continued from Figure 7

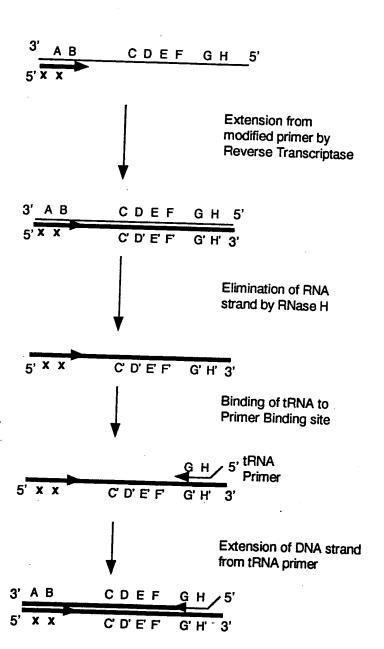


Figure 8

RNA with Ligands on Primer (Continued)

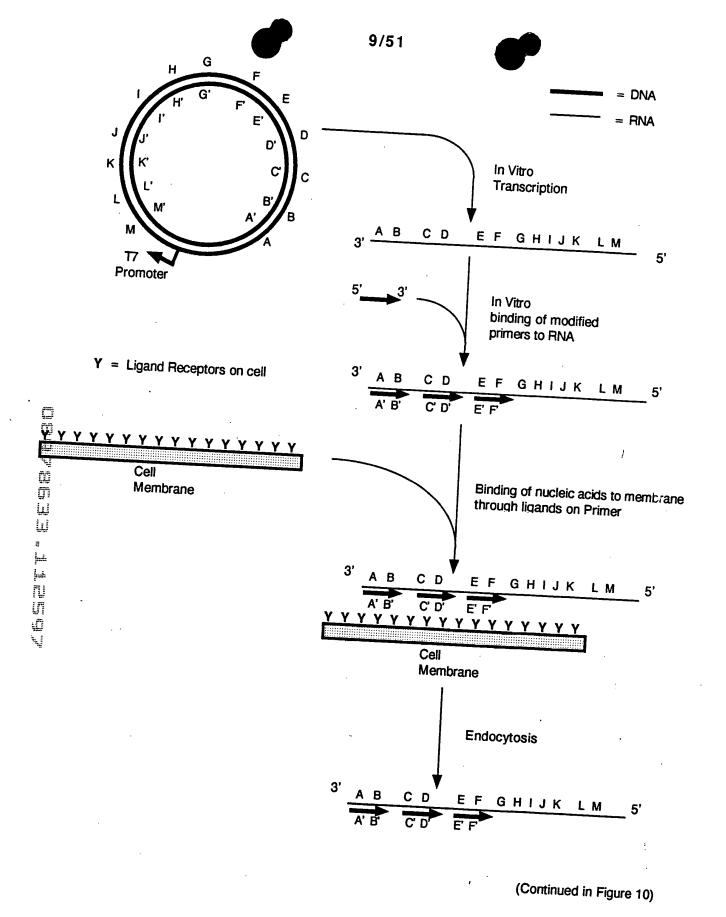


Figure 9
RNA with Ligands on Multiple Primers



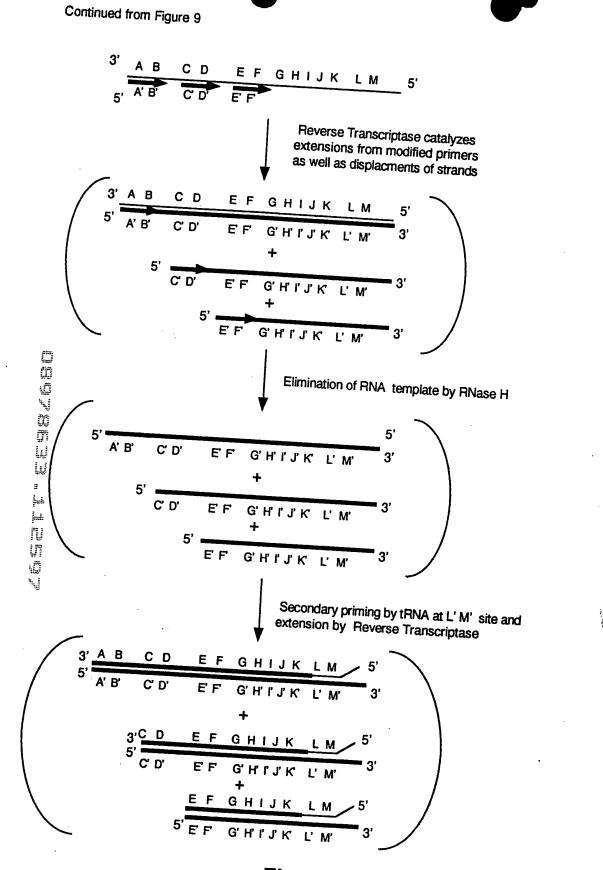


Figure 10 RNA with Ligands on Multiple Primers (Continued)

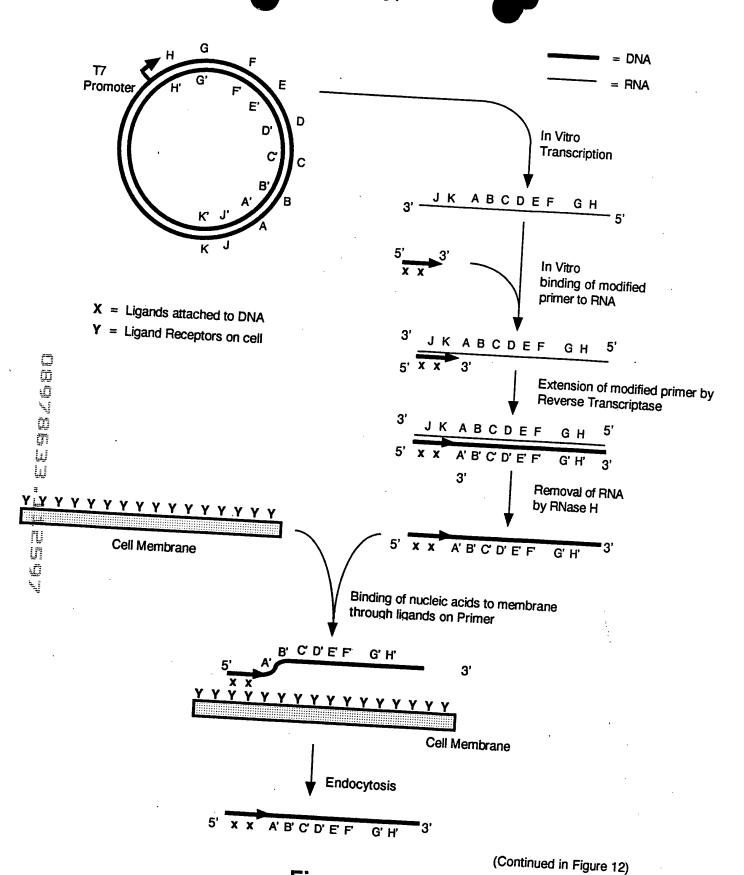
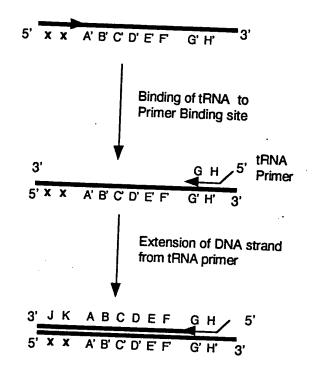
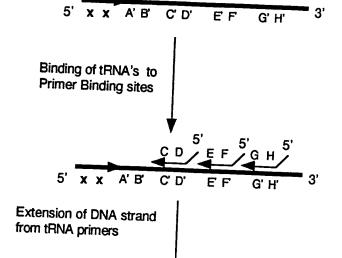


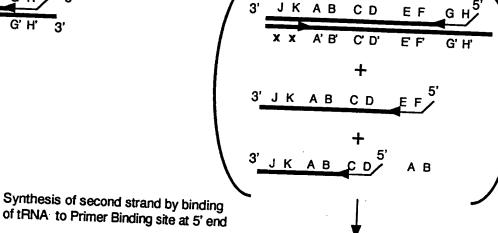
Figure 11
Single-stranded DNA with attached Ligands

(a) Presence of a single tRNA primer site



(b) Presence of multiple tRNA primer sites





E F

Figure 12

Single-stranded DNA with attached Ligands (continued)

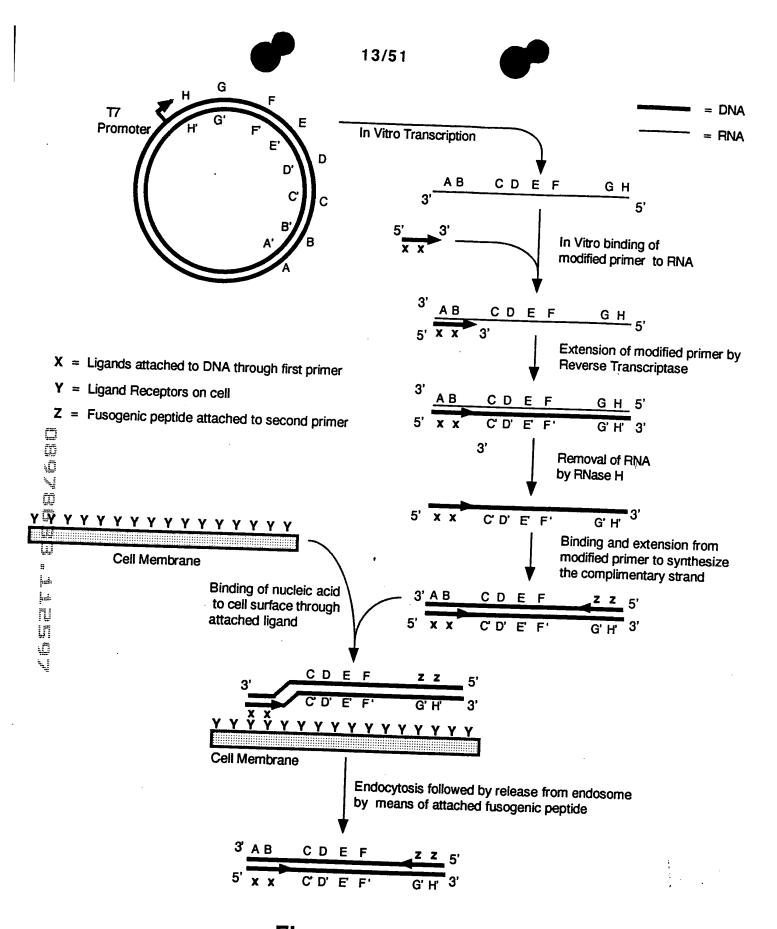


Figure 13
Linear Double-stranded DNA with attached Moieties on each strand

Figure 14
Enhanced Delivery of Retroviral Vector to Haematopoeitic Stem Cell

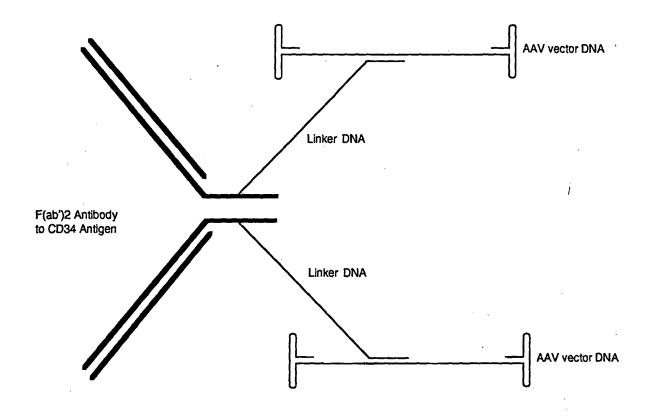


Figure 15
Enhanced Delivery of Vector
DNA to Haematopoeitic Stem Cell

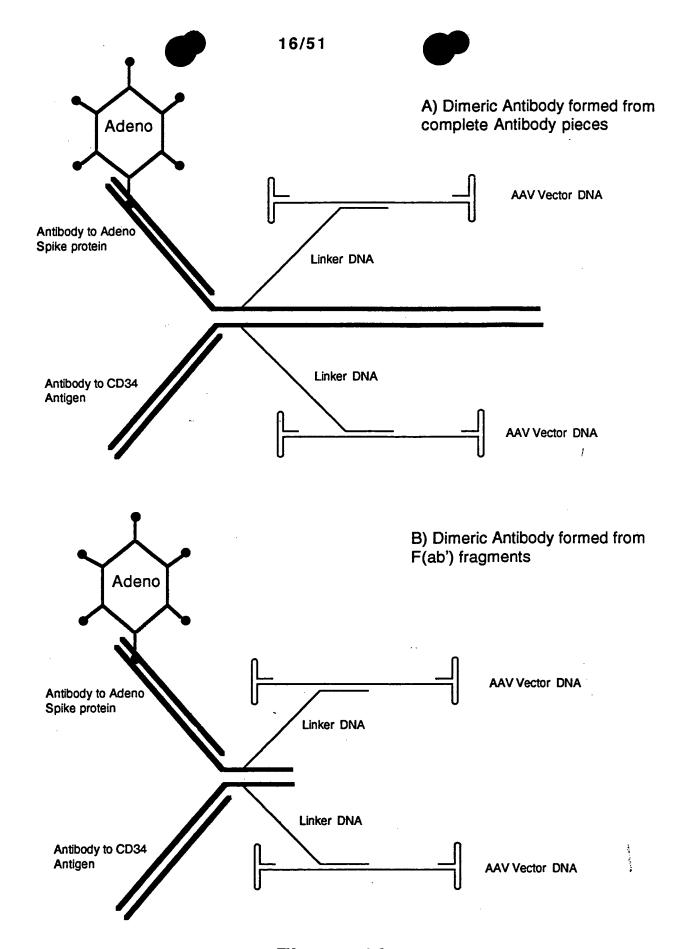


Figure 16
Covalent Attachment of vector DNA to Dimeric Antibody

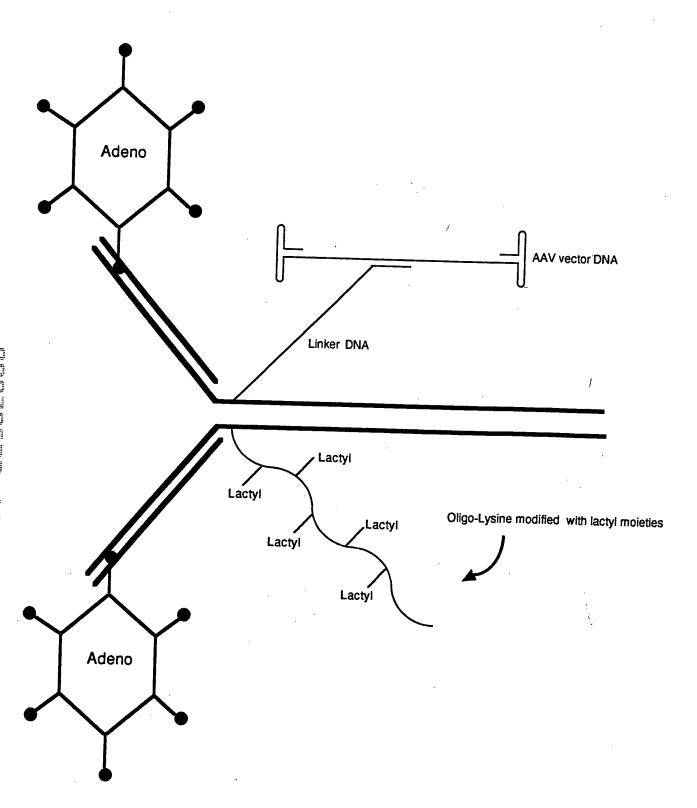


Figure 17
Covalent attachment of Modified DNA to a Monovalent Antibody

Figure 18
Modified DNA used as a Binder

OBSYSSELL LIESSy



NH₂

IV

(continued in Figure 20)

Figur 19 Synthetic Steps for Creation of Antibodies With Nucleic Acid Moieties Attached





Figure 20 **Continuation of Synthetic Steps**

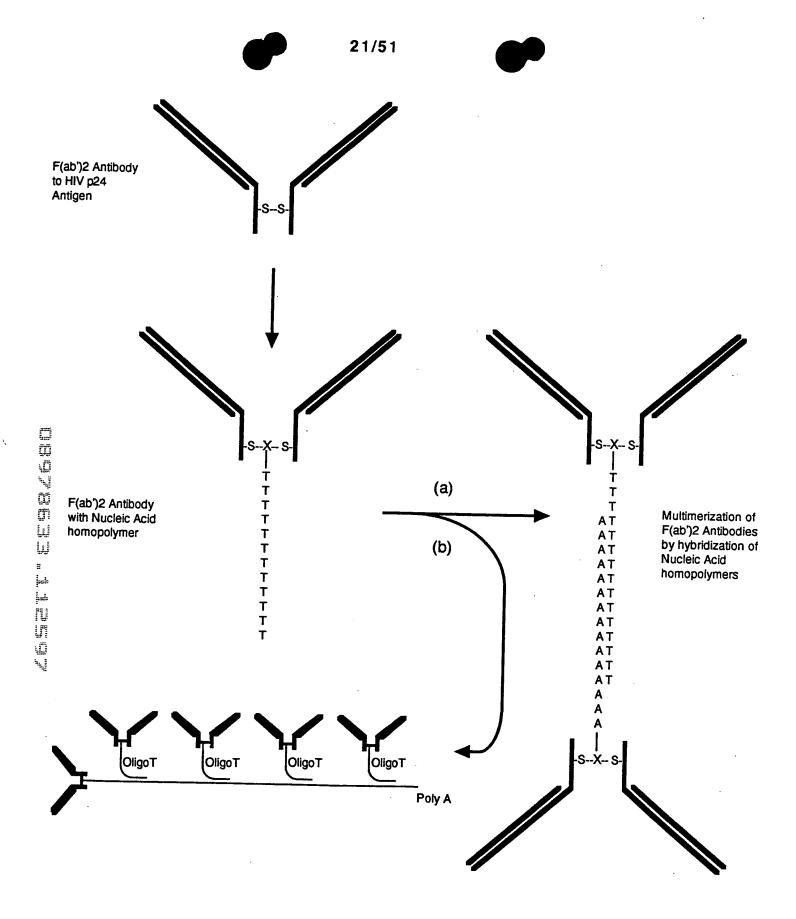


Figure 21
Enhanced Binding of Antibodies to Antigens by Multimerization

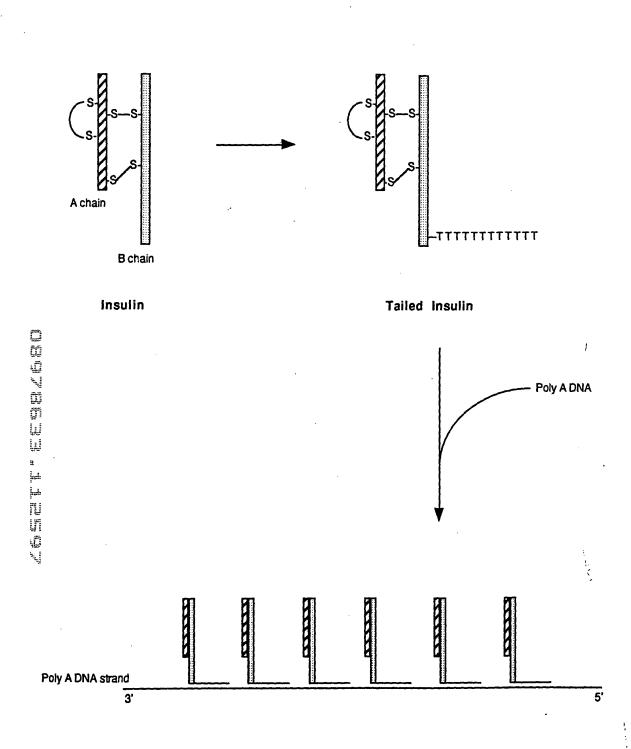


Figure 22
High Affinity Multi-Insulin Soluble Complex

Multimerization of Insulin molecules by hybridization to discrete Sequences

Figure 23

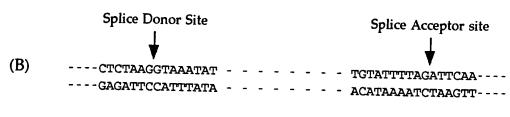




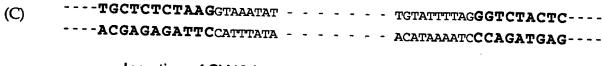
Intron insertion site

(A) ----TGCTCTCTAAGGGTCTACTC----

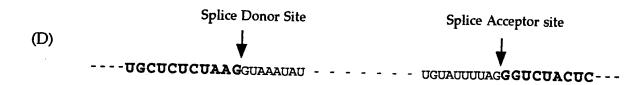
T7 RNA Polymerase Sequence



SV40 Intron Sequence



Insertion of SV40 Intron into polymerase coding sequence



mRNA transcript containing intron

(E)
----UGCUCUCUAAGGGUCUACUC--mRNA transcript after splicing has normal T7 Sequence

Figure 24

Fusion of Intron into T7 RNA Polymerase Coding Sequence

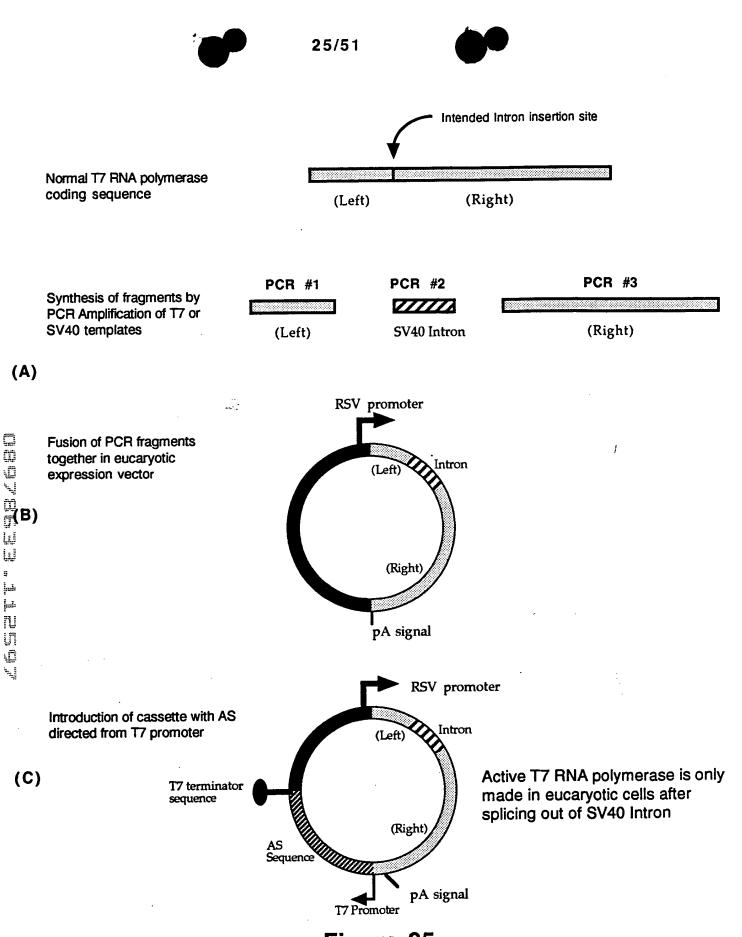
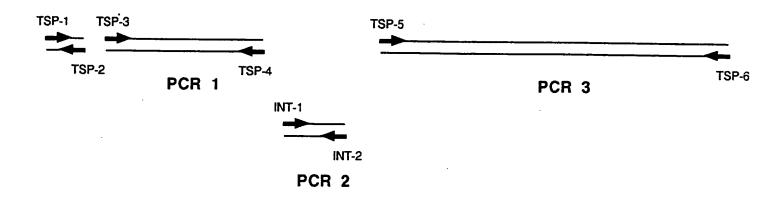


Figure 25
Construction of T7 Expression Vector

A) Synthesis of pieces



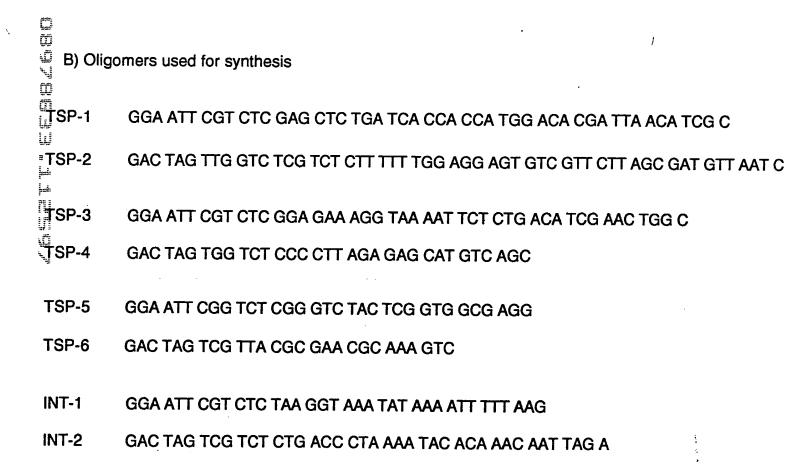


Figure 26

Synthesis of Pieces for Construction of T7 RNA Polymerase with Intron





Formation of Nuclear Localisation Signal by Fusion of TSP1/TSP2 Product to Clone with PCR #1 product

Annealing of TSP1 with TSP2

1517.1 5' gg aat teg tet ega get etg ate aec aec aig gae aeg att aac ate ge 3' 3' e taa tig tag ega tie tig ega gaa gga tit tit etg ega get tit tee tet get etg git gat eag s' TSP2

Extension of TSP1/TSP2 by polymerase

5' GG AAT TCG TCT CGA GCT CTG ATC ACC ACC ATG GAC ACG ATT AAC ATC GCT AAG AAC GAC ACT CCT CCA AAA AAG AGA CGA GAC CAA CTA GTC 3' CC TTA AGC AGA GCT CGA GAC GTA TGG TGG TAC CTG TGG TAA TTG TAG CGA TTC TTG TGA GGA GGT TTT TTC TCT G<u>CT CTG G</u>TT GAT CAG

Digestion of TSP1/TSP2 product with Bsa I

Digestion of PCR #1 clone (pL-1) with BsmB

5' GGA AITT COT CTC G Bem Bl

CCT TAA GCA GAG CCTCT

Ligation of Bsa I digested TS1/TS2 product to BsmB I digested PCR#1 clone

5' GG AAT TCG TCT CGA GCT CTG ATC ACC ACC ATG GAC ACG ATT AAC ATC GCT AAG AAC GAC ACT CCT CCA AAA AAG AAA AAG 3' CC TTA AGC AGA GCT CGA GAC GTA TGG TGG TAC CTG TGC TAA TTG TAG CGA TTC TTG-CTG TGA GGA GGT TTT TTC TCT TTG

MAG

GAC CG-----TCT GAC ATC GAA TAG · CTT CIO AGA

Figure

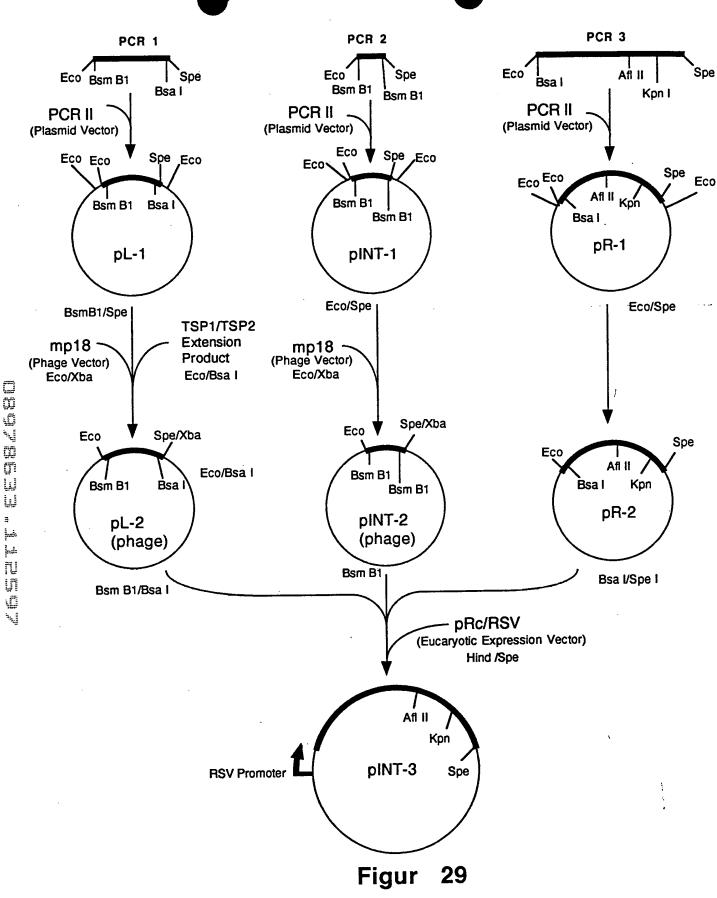
Comparison of the 5' ends of the Nucleotide Sequences of Wild Type and Modified T7 RNA Polymerase

Wild Type T7 nucleic and amino acid sequence

Modified T7 nucleic and amino acid sequence with Nuclear Localisation Signal (NLS) insertion

RITO GAC ACG AITT AAC AITC GCT AAG AAC GAC ACT CCT CCA AAA AAG AGA AAG GTA AAA TIC TCT GAC AIC GAA CTG GC----GAC CG---TGA GGA GGT TITT TITC TOT TITC CAT TITT AND AGA OTO THE CIT TAC CTG TGC TAA TTG TAG CGA TTC TTG CTG 01 6

Figure 28



Fusion of PCR Pieces to Construct T7 RNA Polymerase with an Intron





GAT CAT TAG ACC AGA TCT GAG CCT GGG AGC TCT CTG GCT AAC TAG GGA ACC CAC TGCTTA AGC CTC AAG HTA-1 GAT CCT TGA GGC TTA AGC AGT GGG TTC CCT AGT TAG CCA GAG AGC TCC CAG GCT CAG ATC TGG TCT AAT HTA-2 HTB-1 GAT CAC CTT AGG CTC TCC TAT GGC AGG AAG AAG CGG AGA CAG CGA AGA CCT CCT CAA G GAT CCT TGA GGA GGT CTT CGT CGC TGT CTC CGC TTC TTC CTG CCA TAG GAG AGC CTA AGG T HTB-2 GAT CAT AGT GAA TAG AGT TAG GCA GGG ATA CTC ACC ATT ATC GTT TCA GAC CCA CCT CCC AG HTC-1 GAT CCT GGG AGG TGG GTC TGA AAC GAT AAT GGT GAG TAT CCC TGC CTA ACT CTA TTC ACT AT HTC-2 TER-1 AAT CTA GAG CTA ACA AAG CCC GAA AGG AAG TER-2 TTC TGC AGA TAT AGT TCC TCC TTT CAG C

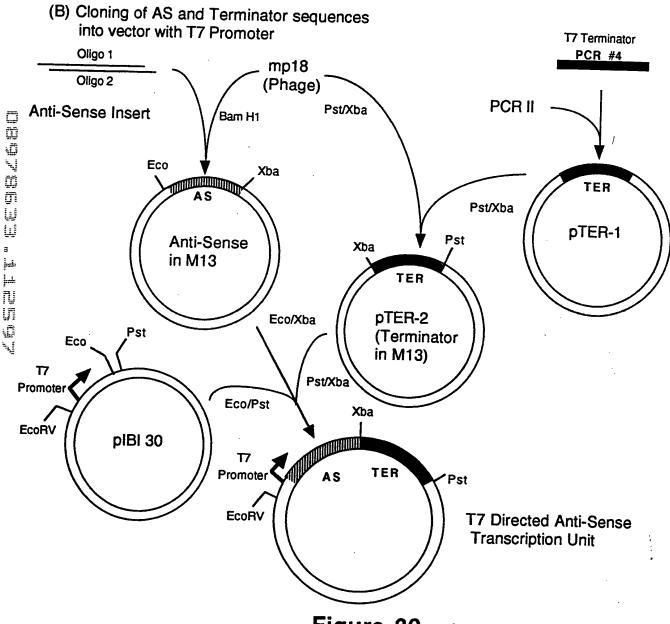


Figure 30 Insertion of Anti-Sense Sequences into T7 Directed Transcription Units

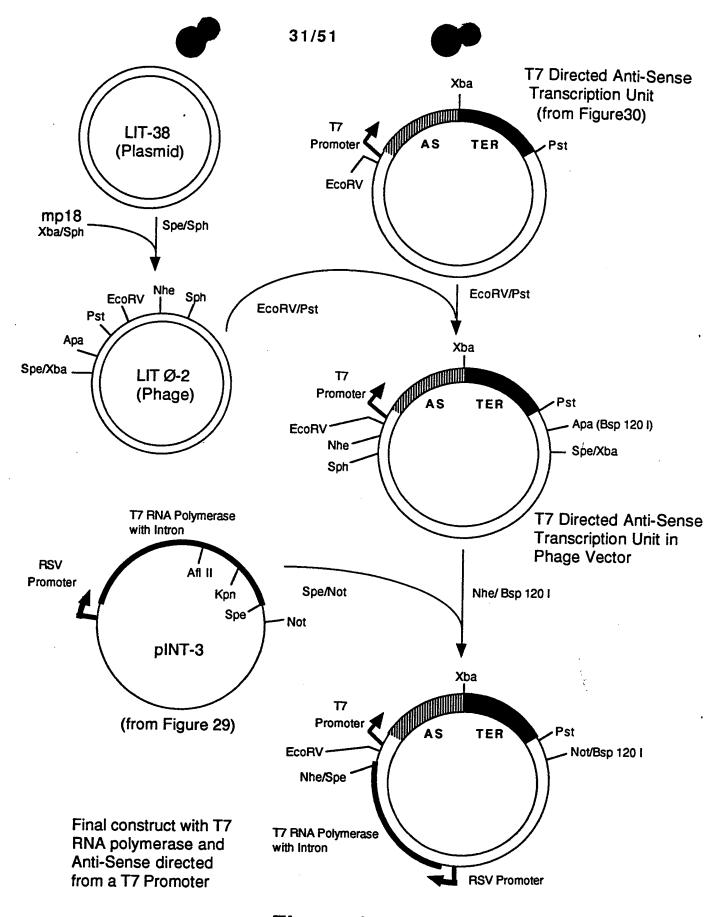


Figure 31
Construct with T7 RNA polymerase and Anti-Sense directed from a T7 Promoter





A) Oligomers for introduction of T7 signals and polylinker

- PL-1

 TCG AGC CAT GGC TTA AGG ATC CGT ACG TCC GGA GCT AGC GGG CCC ATC GAT ACT

 AGT TAA ATG CAG ATC T
- PL-2 CTA GAG ATC TGC ATT TAA CTA GTA TCG ATG GGC CCG CTA GCT CCG GAC GTA CGG
 ATC CTT AAG CCA TGG C

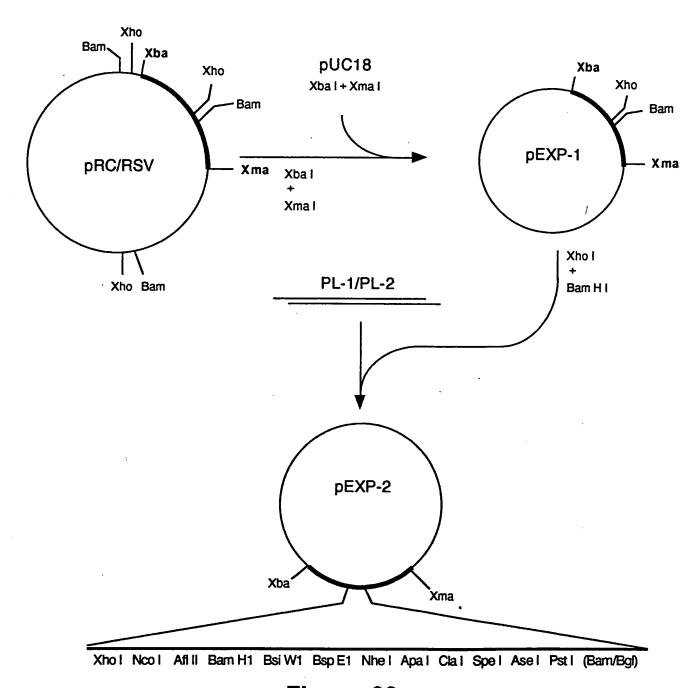
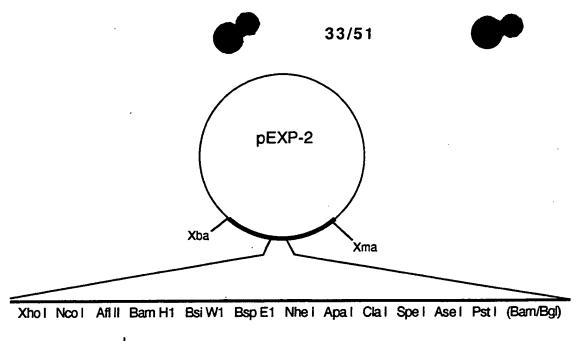


Figure 32

Introduction of Poly-Linker for Creation of Protein Expression Vector



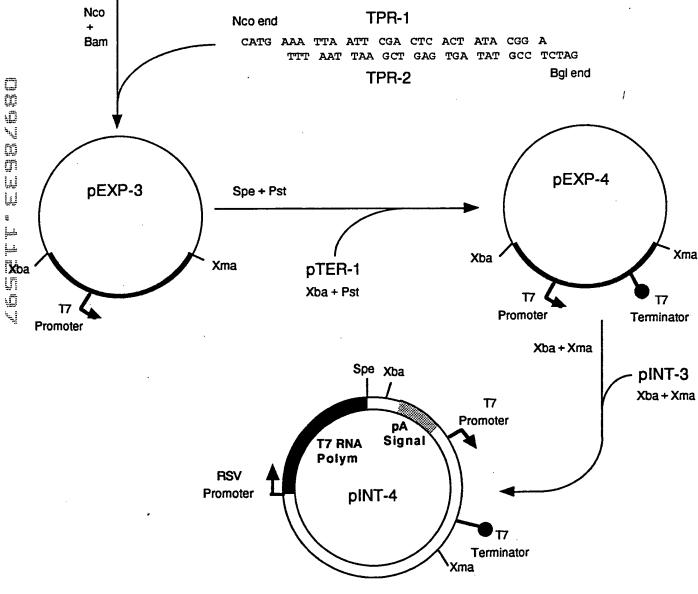
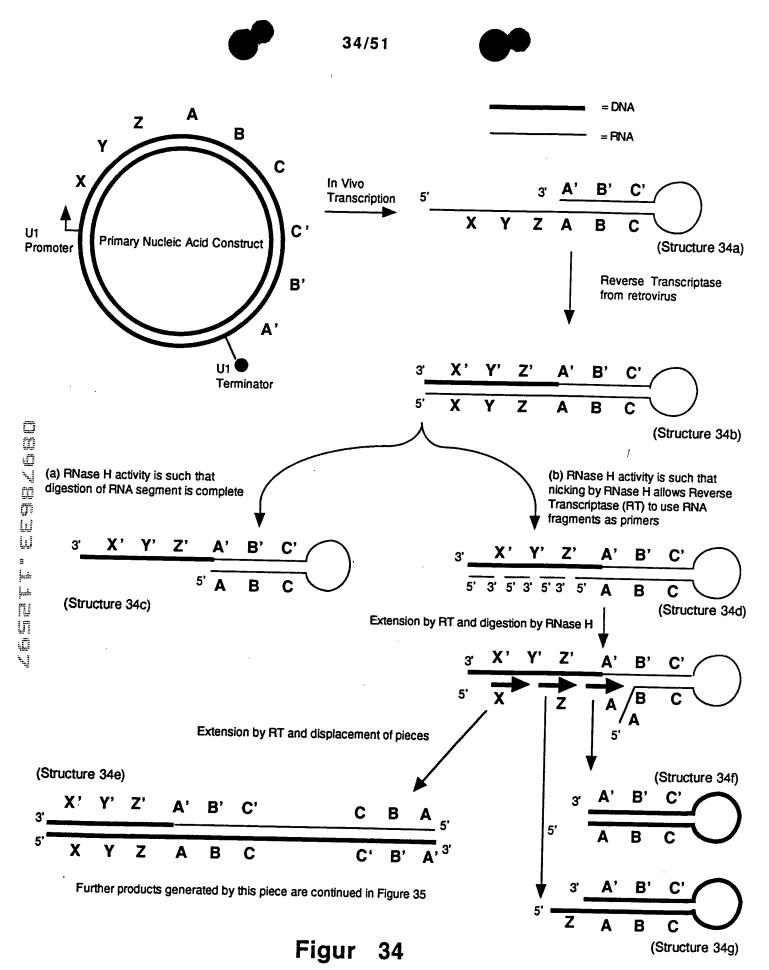


Figure 33
Final steps for construction of Expression Vector

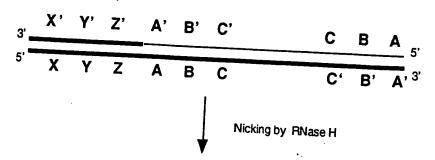


Construct that produces single-stranded Anti-Sense DNA

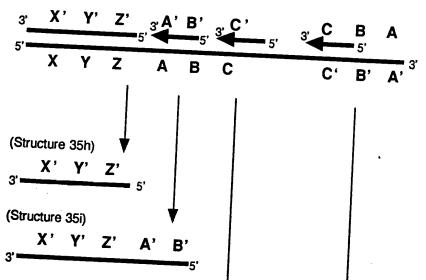




(Structure 34e)



Extension by RT and digestion by RNase H



(Structure 35j)

X' Y' Z' A' B' C'

Extension by RT and displacement generates Single-Stranded DNA and a mostly Double-stranded DNA molecule

(Structure 35k)

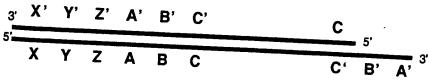


Figure 35
Continuation of Process from Figure 34

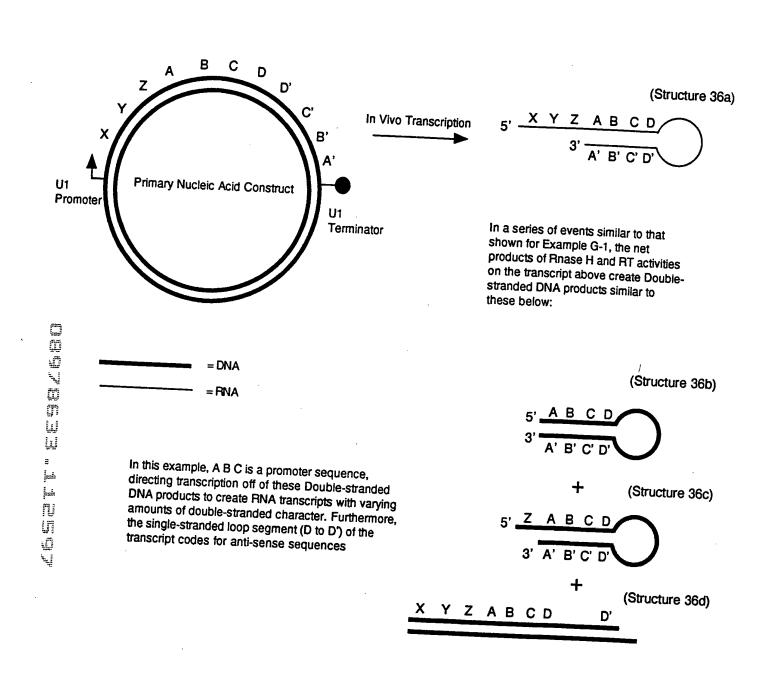


Figure 36

Construct that produces RNA that is Reverse Transcribed to create Secondary DNA Constructs capable of directing transcription

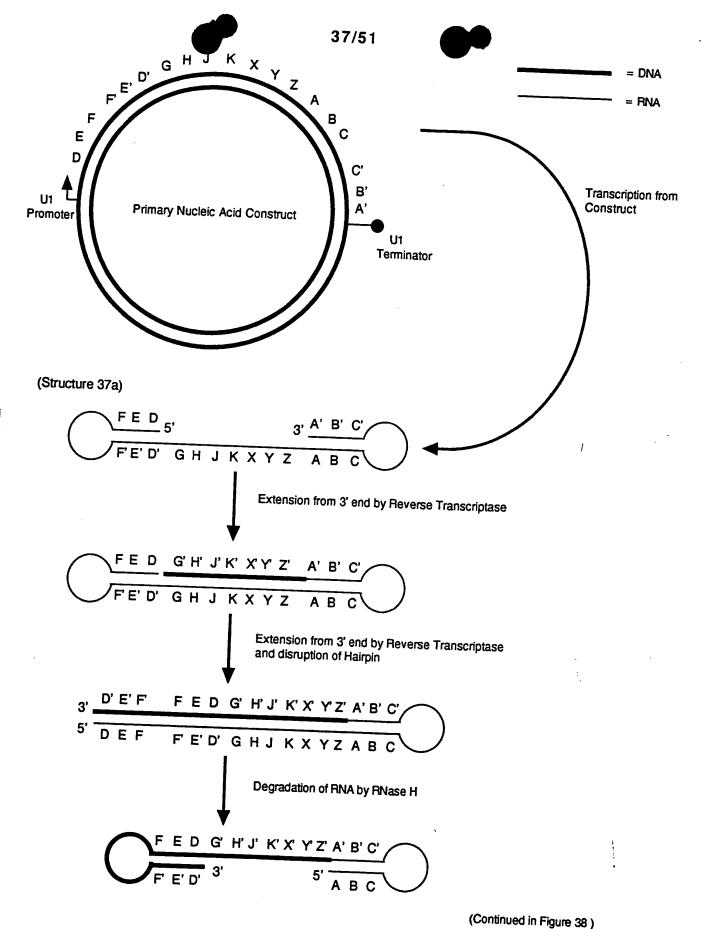
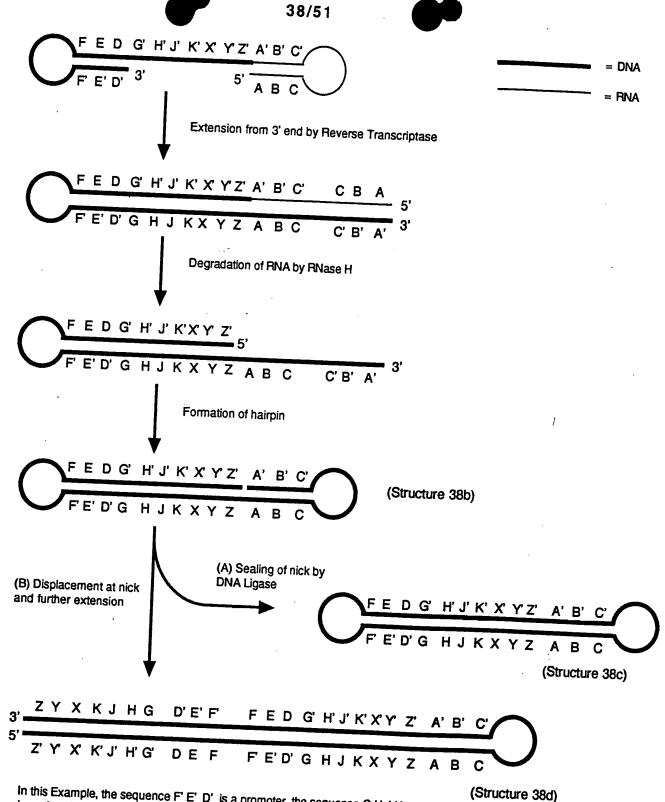


Figure 37
Construct which Propagates a Double Hairpin Production Center



In this Example, the sequence F' E' D' is a promoter, the sequence G H J κ is an Anti-Sense sequence and X Y Z is a Poly A signal

Figure 38 Continuation of process from Figure 37

Figure 39

Construct which propagates a Production Center capable of Inducible Suicide

Figure 40
Use of tRNA primers to create a DNA construct for secondary production of transcripts

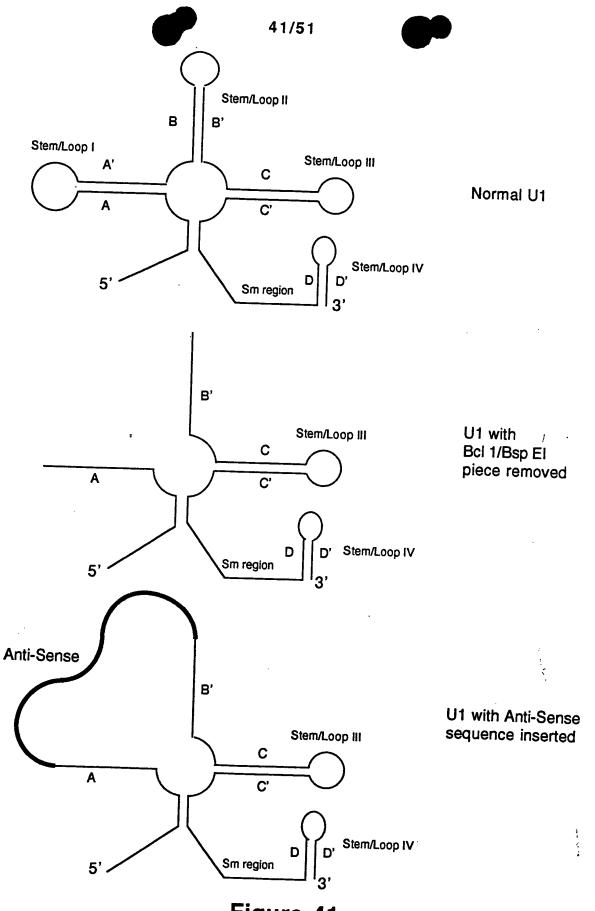


Figure 41
Excision of Sequences from U1 Transcript Region and Replacement with Novel Sequences





(A) Anti-sense oligomers

HVA-1

GAT CCG GAT TGA GGC TTA AGC AGT GGG TTC CCT AGT TAG CCA GAG AGC TCC CAG GCT CAG ATC TGG TCT AAT

HVA-2

CCG GAT TAG ACC AGA TCT GAG CCT GGG AGC TCT CTG GCT AAC TAG GGA ACC CAC TGC TTA AGC CTC AAT CCG

HVB-1

GAT CCG GAC CTT GAG GAG GTC TTC GTC GCT GTC TCC GCT TCT TCC TGC CAT AGG AGA GCC TAA GGT

HVB-2

CCG GAC CTT AGG CTC TCC TAT GGC AGG AAG AAG CGG AGA CAG CGA AGA CCT CCT CAA GGT CCG

HVC-1

GAT CCG GAT GGG AGG TGG GTC TGA AAC GAT AAT GGT GAG TAT CCC TGC CTA ACT CTA TTC ACT AT

HVC-2

CCG GAT AGT GAA TAG AGT TAG GCA GGG ATA CTC ACC ATT ATC GTT TCA GAC CCA CCT CCC ATC CG

HVD-1

GAT CAG CAT GCC TGC AGG TCG ACT CTA GAC CCG GGT ACC GAG CTC GCC CTA TAG TGA GT C GTA TTA T

HVD-2

CCG GAT AAT ACG ACT CAC TAT AGG GCG AGC TCG GTA CCC GGG TCT AGA GTC GAC CTG CAG GCA TGC T

(B) Replacement of U1 sequences with HIV Anti-sense sequences

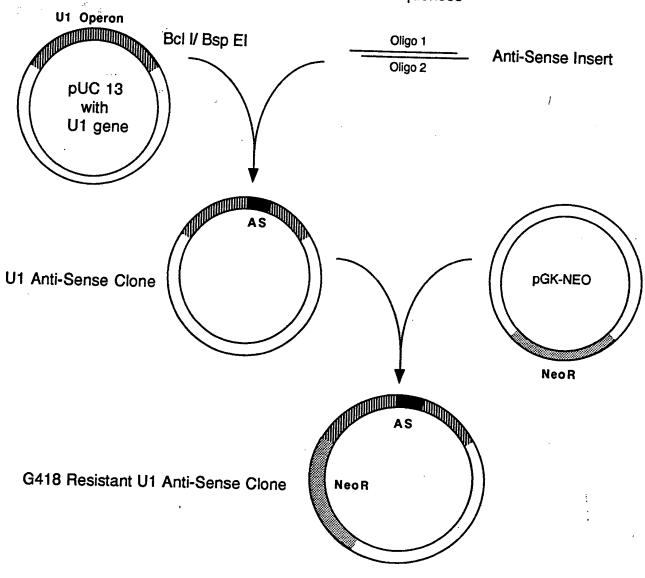


Figure 42
Insertion of Anti-Sense Sequences into U1Operons

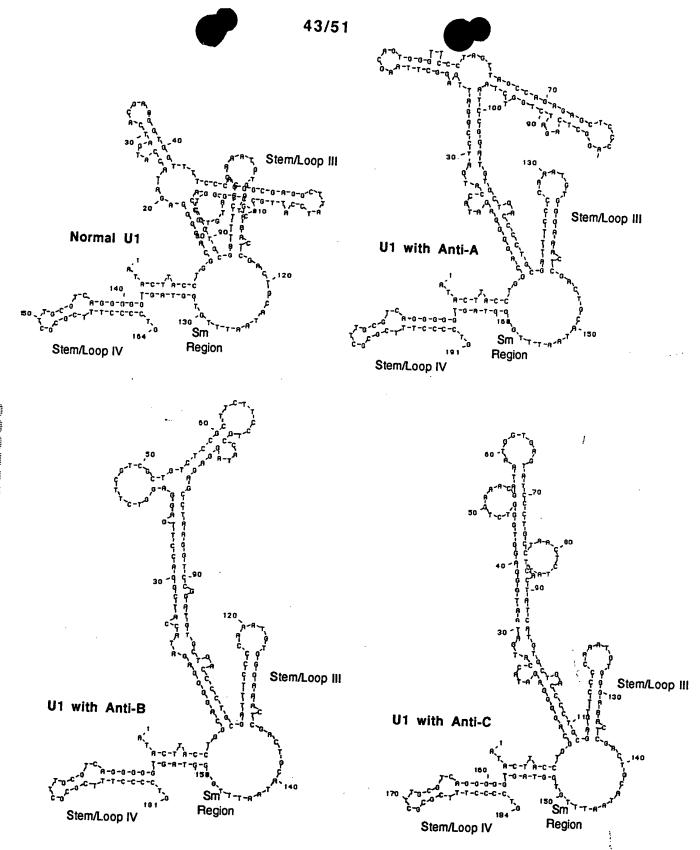


Figure 43
Predicted Secondary structures for U1
Transcripts with Anti-sense Substitutions

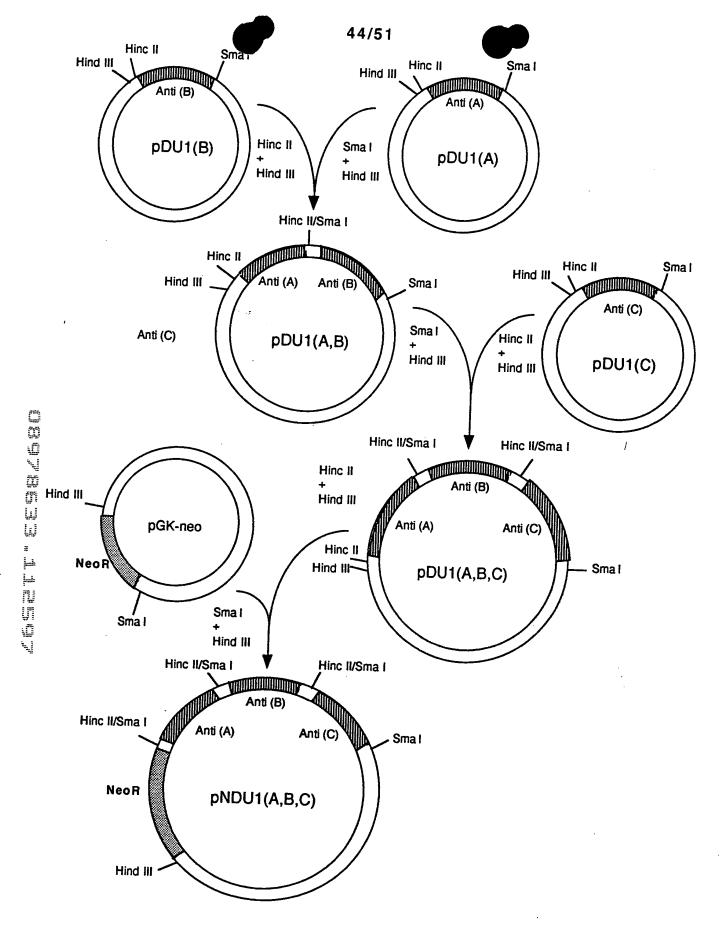


Figure 44
Construction of U1 Multiple Operon Clone

Figure 45
Construction of T7 Triple Operon

pNDU1(A,B,C)

Triple U1 Operon Construct with HIV Anti-Sense

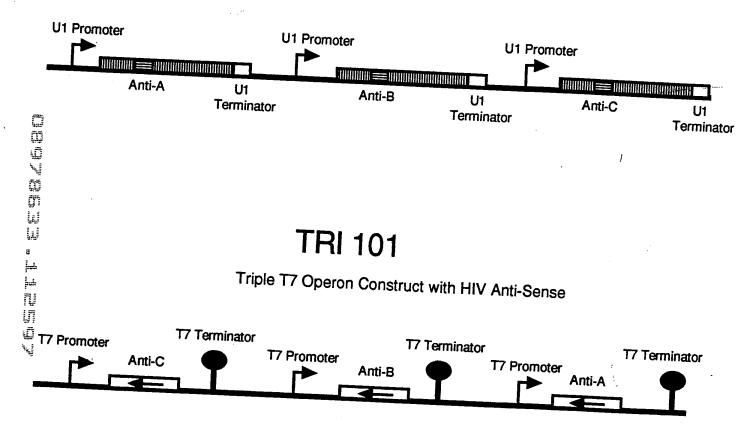


Figure 46
Structures of Triple Operon Constructs from Figures 44 and 45

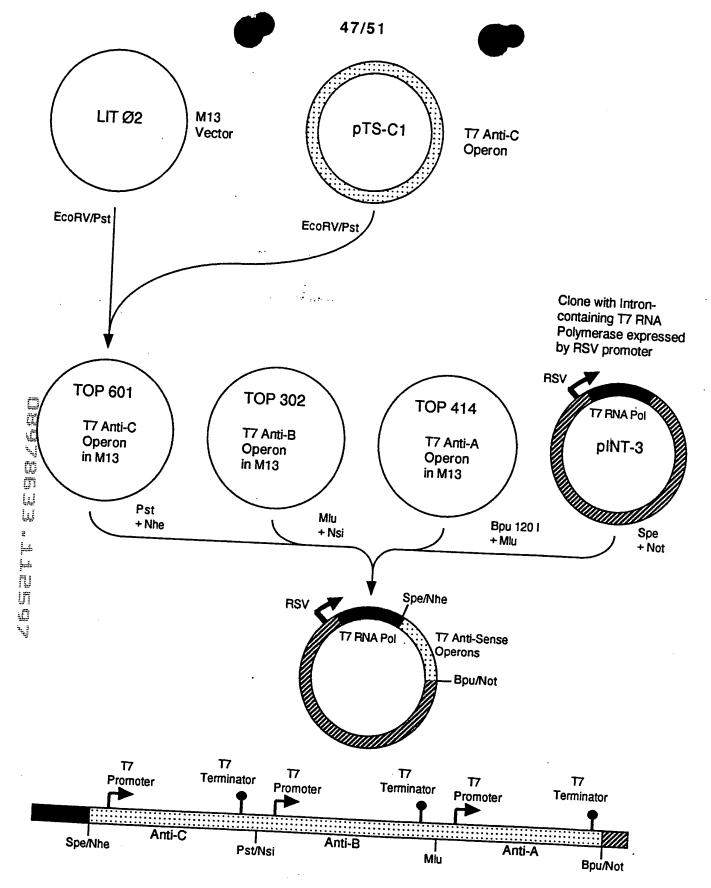
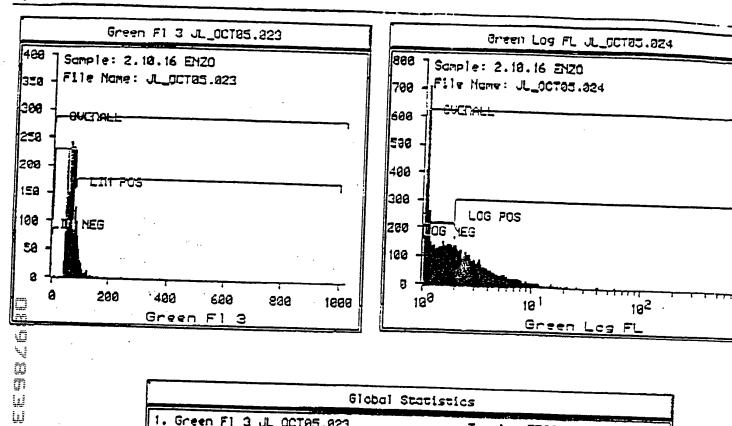


Figure 47
Construction of Multiple T7 Operons in Vector coding for T7 RNA Polymerse



Global Statistics					
1. Green F1 3 JL_OCT05.023 2. Green Log FL JL_OCT05.024 Hist Region Bounds Counts 1. LIN MEG 1 78 5714 LIN POS 85 1002 1129 OVERALL 1 1024 7509 2. LOG MEG 2 2 4211 LOG POS 2 1001 7509 OVERALL 2 1001 7509	Total = 7509 Total = 7509 Total = 7509 X	XC 144 177 231 69 88			

Figure 48

Flow cytometry data measuring binding of anti-CD4+ antibody to HIV resistant U037 cells

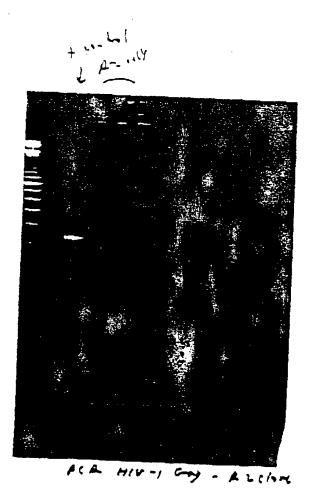


Figure 49

PCR amplification of gag region indicating absence of HIV in viral resistant cell line (2.10.16) after challenge

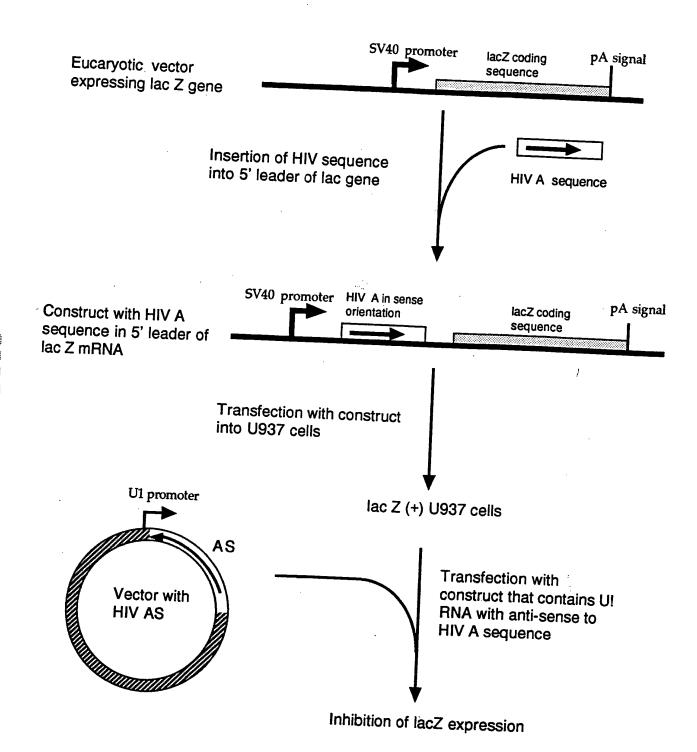


Figure 50

Clone with target-lacZ fusion will have reduced expression of lacZ after transfection by HIV Anti-sense construct





51/51

Enzyme activity as expressed by A_{420} readings in extracts prepared from

	propercy nom		
•••	2.5 x 10 ⁴ cells	5 x 10 ⁴ cells	1.0 x 10 ⁵ cells
U 937 [untransfected]	0.018	0.023	0.034
U 937 [HIV A clone]	0.154	0.277	0.566
U937 [HIV A / Anti-A]	0.010	0.017	0.027
U 937 [HIV A/Anti-ABC]	0.013	0.021	0.035
U 937 [HIV A/Null DNA]	0.120	0.212	0.337

[B] Expression of Beta-galactosidase activity by In situ assay:

U 937 [untransfected] no blue spots in cells

U 937 [HIV A clone] blue spots in cells

U 937 [HIV A/Anti A] no blue spots in cells

U 937 [HIV A/Anti ABC] no blue spots in cells

U 937 [HIV A / Null DNA] blue spots in cells

Figure 51

Expression of Beta-galactosidase activity in extracts